

4. Gaylor  
PAGE: 1

1642

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/776,044

DATE: 02/11/98  
TIME: 09:51:30

INPUT SET: S23406.raw

This Raw Listing contains the General  
Information Section and up to the first 5 pages.

ENTERED

SEQUENCE LISTING

1  
2  
3 (1) General Information:  
4  
5 (i) APPLICANT: BYWATER, MARGARET  
6 LINDSTROM, PER  
7 INGANAS, MATS  
8  
9 (ii) TITLE OF INVENTION: SEQUENCE-BASED MUTATION ANALYSIS OF  
10 NEOPLASTIC TISSUE FOR DIAGNOSIS OR PROGNOSIS OF THE  
11 NEOPLASIA  
12  
13 (iii) NUMBER OF SEQUENCES: 25  
14  
15 (iv) CORRESPONDENCE ADDRESS:  
16 (A) ADDRESSEE: BIRCH, STEWART, KOLASCH AND BIRCH  
17 (B) STREET: PO BOX 747  
18 (C) CITY: FALLS CHURCH  
19 (D) STATE: VA  
20 (E) COUNTRY: USA  
21 (F) ZIP: 22040-0747  
22  
23 (v) COMPUTER READABLE FORM:  
24 (A) MEDIUM TYPE: Floppy disk  
25 (B) COMPUTER: IBM PC compatible  
26 (C) OPERATING SYSTEM: PC-DOS/MS-DOS  
27 (D) SOFTWARE: PatentIn Release #1.0, Version #1.30  
28  
29 (vi) CURRENT APPLICATION DATA:  
30 (A) APPLICATION NUMBER:  
31 (B) FILING DATE:  
32 (C) CLASSIFICATION:  
33  
34 (viii) ATTORNEY/AGENT INFORMATION:  
35 (A) NAME: MURPHY JR, GERALD M  
36 (B) REGISTRATION NUMBER: 28,977  
37 (C) REFERENCE/DOCKET NUMBER: 1614-178  
38  
39 (ix) TELECOMMUNICATION INFORMATION:  
40 (A) TELEPHONE: (703) 205-8000  
41 (B) TELEFAX: (703) 205-8050  
42  
43  
44 (2) INFORMATION FOR SEQ ID NO:1:  
45  
46 (i) SEQUENCE CHARACTERISTICS:

RAW SEQUENCE LISTING  
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47 (A) LENGTH: 19 base pairs  
48 (B) TYPE: nucleic acid  
49 (C) STRANDEDNESS: single  
50 (D) TOPOLOGY: linear  
51

52 (ii) MOLECULE TYPE: other nucleic acid  
53 (A) DESCRIPTION: /desc = "PRIMER"  
54  
55  
56

57  
58 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:  
59

60 GTGCTTTCCA CGACGGTG A

19

61  
62 (2) INFORMATION FOR SEQ ID NO:2:  
63

64 (i) SEQUENCE CHARACTERISTICS:  
65 (A) LENGTH: 23 base pairs  
66 (B) TYPE: nucleic acid  
67 (C) STRANDEDNESS: single  
68 (D) TOPOLOGY: linear  
69

70 (ii) MOLECULE TYPE: other nucleic acid  
71 (A) DESCRIPTION: /desc = "PRIMER"  
72  
73  
74  
75

76 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:  
77

78 TCATGTGCTG TGACTGCTTG TAG

23

79  
80 (2) INFORMATION FOR SEQ ID NO:3:  
81

82 (i) SEQUENCE CHARACTERISTICS:  
83 (A) LENGTH: 19 base pairs  
84 (B) TYPE: nucleic acid  
85 (C) STRANDEDNESS: single  
86 (D) TOPOLOGY: linear  
87

88 (ii) MOLECULE TYPE: other nucleic acid  
89 (A) DESCRIPTION: /desc = "PRIMER"  
90  
91  
92  
93

94 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:  
95

96 GGCGGGGGTG TGGAATCAA

19

97  
98 (2) INFORMATION FOR SEQ ID NO:4:  
99

RAW SEQUENCE LISTING  
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100 (i) SEQUENCE CHARACTERISTICS:  
101 (A) LENGTH: 23 base pairs  
102 (B) TYPE: nucleic acid  
103 (C) STRANDEDNESS: single  
104 (D) TOPOLOGY: linear  
105  
106 (ii) MOLECULE TYPE: other nucleic acid  
107 (A) DESCRIPTION: /desc = "PRIMER"  
108  
109  
110  
111

112 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:  
113

114 TCTGGCATTC TGGGAGCTTC ATC  
115

23

116 (2) INFORMATION FOR SEQ ID NO:5:  
117

118 (i) SEQUENCE CHARACTERISTICS:  
119 (A) LENGTH: 20 base pairs  
120 (B) TYPE: nucleic acid  
121 (C) STRANDEDNESS: single  
122 (D) TOPOLOGY: linear  
123

124 (ii) MOLECULE TYPE: other nucleic acid  
125 (A) DESCRIPTION: /desc = "PRIMER"  
126  
127  
128  
129

130 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:  
131

132 CCGTCCCAGT AGATTACCAC  
133

20

134 (2) INFORMATION FOR SEQ ID NO:6:  
135

136 (i) SEQUENCE CHARACTERISTICS:  
137 (A) LENGTH: 20 base pairs  
138 (B) TYPE: nucleic acid  
139 (C) STRANDEDNESS: single  
140 (D) TOPOLOGY: linear  
141

142 (ii) MOLECULE TYPE: other nucleic acid  
143 (A) DESCRIPTION: /desc = "PRIMER"  
144  
145  
146  
147

148 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:  
149

150 GTTTTCCGTC TGGGCTTCTT  
151

20

152 (2) INFORMATION FOR SEQ ID NO:7:

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153  
154 (i) SEQUENCE CHARACTERISTICS:  
155 (A) LENGTH: 27 base pairs  
156 (B) TYPE: nucleic acid  
157 (C) STRANDEDNESS: single  
158 (D) TOPOLOGY: linear  
159  
160 (ii) MOLECULE TYPE: other nucleic acid  
161 (A) DESCRIPTION: /desc = "PRIMER"  
162  
163  
164  
165  
166 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:  
167  
168 CTGTGACTTG CACGTACTCC CCTGCCC 27  
169  
170 (2) INFORMATION FOR SEQ ID NO:8:  
171  
172 (i) SEQUENCE CHARACTERISTICS:  
173 (A) LENGTH: 23 base pairs  
174 (B) TYPE: nucleic acid  
175 (C) STRANDEDNESS: single  
176 (D) TOPOLOGY: linear  
177  
178 (ii) MOLECULE TYPE: other nucleic acid  
179 (A) DESCRIPTION: /desc = "PRIMER"  
180  
181  
182  
183  
184 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:  
185  
186 TAGACTGACC CTTTTTGGAC TTC 23  
187  
188 (2) INFORMATION FOR SEQ ID NO:9:  
189  
190 (i) SEQUENCE CHARACTERISTICS:  
191 (A) LENGTH: 21 base pairs  
192 (B) TYPE: nucleic acid  
193 (C) STRANDEDNESS: single  
194 (D) TOPOLOGY: linear  
195  
196 (ii) MOLECULE TYPE: other nucleic acid  
197 (A) DESCRIPTION: /desc = "PRIMER"  
198  
199  
200  
201  
202 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:  
203  
204 CGTGTGGAGT ATTTGGATGA C 21  
205

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206 (2) INFORMATION FOR SEQ ID NO:10:

207

208 (i) SEQUENCE CHARACTERISTICS:

209 (A) LENGTH: 22 base pairs

210 (B) TYPE: nucleic acid

211 (C) STRANDEDNESS: single

212 (D) TOPOLOGY: linear

213

214 (ii) MOLECULE TYPE: other nucleic acid

215 (A) DESCRIPTION: /desc = "PRIMER"

216

217

218

219

220 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

221

222 TGGTTTCTTC TTTGGCTGGG GA

22

223

224 (2) INFORMATION FOR SEQ ID NO:11:

225

226 (i) SEQUENCE CHARACTERISTICS:

227 (A) LENGTH: 21 base pairs

228 (B) TYPE: nucleic acid

229 (C) STRANDEDNESS: single

230 (D) TOPOLOGY: linear

231

232 (ii) MOLECULE TYPE: other nucleic acid

233 (A) DESCRIPTION: /desc = "PRIMER"

234

235

236

237

238 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

239

240 GCTTTGAGGT GCGTGTTTGT G

21

241

242 (2) INFORMATION FOR SEQ ID NO:12:

243

244 (i) SEQUENCE CHARACTERISTICS:

245 (A) LENGTH: 21 base pairs

246 (B) TYPE: nucleic acid

247 (C) STRANDEDNESS: single

248 (D) TOPOLOGY: linear

249

250 (ii) MOLECULE TYPE: other nucleic acid

251 (A) DESCRIPTION: /desc = "PRIMER"

252

253

254

255

256 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:

257

258 CTGTCACTGG GGAACAAGAA G

21

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**SEQUENCE VERIFICATION REPORT**  
**PATENT APPLICATION US/08/776,044**

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***INPUT SET: S23406.raw***

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